SEQUENCE LISTING

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<110> Williams, Richard B.
 <120> SYSTEM AND METHODS FOR NUCLEIC ACID AND
   POLYPEPTIDE SELECTION
 <130> PRONOV.001BPC
 <150> 09/859,809
 <151> 2001-05-17
<150> 60/206,016
<151> 2000-05-19
 <150> 60/346,965
 <151> 2001-11-16
 <150> 60/529,331
 <151> 2003-12-12
: <150> PCT/US02/37103
 <151> 2002-11-18
 <150> 10/847,087
 <151> 2004-05-17
 <150> 60/625,707
 <151> 2004-11-05
 <150> 10/847,484
 <151> 2004-05-17
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<222> (6)...(6)
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cuagancugg agg
                                                                     13
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 cuagancugg agg
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 <223> n=g, a, u, or c
 <221> misc_feature
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nnnnnccucc agaucuagnn nnn
                                                                      23
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<222> (1) ... (5)
<223> n=g, a, u, or c
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<223> n=g, a, u, or c
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   nnnnnccucc agaucuagnn nnn
                                                                        23
   <210> 5
   <211> 13
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        RNA fragment
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  <223> n=p
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  <223> psoralen bound to G
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 cuagancugg agg
                                                                       13
 <210> 6.
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 <223> chemically synthesized fragment 3; modified tRNA;
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 <223> n=p
<221> misc_feature
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uccugugtnc gauccacaga auucgcaccn
                                                                     30
<210> 7
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<212> DNA
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<222> (22)...(22)
<223> n=p
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   <223> n=puromycin
   <221> modified base
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                                                                         43
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  gcggauuuag cucaguuggg agagcgccag acu
                                                                         33
  <210> 9
  <211> 76.
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        modified tRNA; thymine at residue 54 before
        pseudouridine
 <221> modified_base
 <222> (39)...(39)
 <223> n=p
 <221> modified_base
 \langle 222 \rangle (55) \dots (\overline{5}5)
 <223> n=p
 <221> misc_feature
 <222> (76) ... (76)
 <223> n=puromycin
 <221> misc feature
 <222> (35) ... (36)
 <223> psoralen binding at UA position
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 gcggauuuag cucaguuggg agagcgccag acucuaganc uggagguccu gugtncgauc 60
 cacagaauuc gcaccn
                                                                        76
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                                                                        32
    <210> 11
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  <223> N=p
  <400> 11
  ncuaacnc
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<222> (4)...(4)
<223> n=p
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 \langle 222 \rangle (11)...(12)
 <223> n=p
 <400> 13
 cccnccagag nnagaccc
                                                                            18
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<400> 14
ucuaagneng gagg
                                                                            14
<210> 15
<211> 73
<212> DNA
<213> Artificial Sequence
<223> chemically synthesized; mixed DNA/RNA; alternate
       psoralentated Fragment 1 + 2 + 3
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<223> N-3-methyl uridine
<221> misc_feature
\langle 222 \rangle (36) \dots (37)
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<222> (40)...(40)
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<221> modified base
\langle 222 \rangle (42) \dots (\overline{4}2)
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<221> misc_feature
<222> (73)...(73)
<223> N=puromycin
<400> 15
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 ccacagaauu cgn
 <210> 16
 <211> 32
 <212> RNA
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 <223> Chemically synthesized fragment 1; 3' hydroxyl at
       terminus
 <400> 16
 gcggauuuag cucaguuggg agagcgccag ac
                                                                        32
 <210> 1.7
 <211> 8
 <212> RNA
 <213> Artificial Sequence
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 <223> Chemically synthesized fragment 2; 3' hydroxyl at
       terminus
 <221> modified base
 <222> (1)...(1)
 <223> n=p
<221> modified base
<222> (7)...(7)
<223> n=p
<400> 17
ncuaaanc
                                                                       8
<210> 18
<211> 36
<212> DNA
<213> Artificial Sequence
<223> Chemically synthesized; mixed DNA/RNA fragment 3
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\langle 222 \rangle (15)...(\overline{1}5)
<223> N=p
<221> misc_feature
<222> (36) ... (36)
<223> N=puromycin
<400> 18
uggagguccu gugtncgauc cacagaauuc gcaccn
                                                                       36
<210> 19
<211> 21
<212> RNA
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<221> modified_base <222> (9)(11) <223> N=p	
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<210> 20 <211> 26 <212> RNA <213> Artificial Sequence	
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<400> 20 auauauau auauauau gggggg	26
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220> 223> Chemically synthesized; seq 1B1	

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  <223> no phosphorylation on 5' end
  <400> 23
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                                                                      31
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 <210> 25
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<223> hydroxylated
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<222> (35) ... (36)
<223> crosslinker between residues 35 and 36
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 <211> 55
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 <213> Artificial Sequence
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                                                                     55
 <210> 28
 <211> 61
 <212> RNA
 <213> Artificial Sequence
<220>
<223> Chemically synthesized RNA sequence
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 <222> (59)...(60)
 <223> crosslink between residue 59 and 60
 <221> misc_feature
 <222> (61) ... (61)
<223> g bound to biotin
<221> misc_feature
<222> (33) ... (34)
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<400> 28
ggguuaacuu uagaaggagg ucgccaccau ggnnaaaaug aaaaugaaaa ugaaaaugua 60
g
<210> 29
<211> 61
<212> RNA
<213> Artificial Sequence
<223> Chemically synthesized RNA sequence
<400> 29
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<210> 30
<211> 21
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  <221> modified base
  \langle 222 \rangle (9) \dots (1\overline{1})
  <223> n=p
  <221> misc_feature
 <222> (15)...(21)
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 nnnnnngann nagannnnnn n
                                                                         21
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 <211>.36.
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 <222> (14) ... (14)
 <223> thymine
<221> modified_base
<222> (15)...(15)
<223> n=p
<221> misc_feature
<222> (36) ... (36)
<223> n=puromycin
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uggagguccu gugtnegauc cacagaauuc gcacen
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<210> 32
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  <223> phosphorylated
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  <222> (14)...(14)
  <223> thymine
  <221> modified base
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  <223> n=p
  <221> misc_feature
  <222> (36) ... (36)
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  uggagguccu gugtnegauc cacagaaucu ccacca
                                                               36
  <210> 33
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 <223> mixed DNA/RNA chemically synthesized primer for
 SARS-CoV genome sequence
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                                                           . .
 <222> (32)...(32)
 <223> N-3-methyl uridine
 <221> misc_feature
 <222> (35)...(36)
 <223> psoralen bound to UA
 <221> modified base
 <222> (39)...(39)
 <223> p
 <221> modified base
 \langle 222 \rangle \ (41) \dots (\overline{4}1)
 <223> p
 <221> misc feature
 <222> (72)...(72)
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 geggauuuag eucaguuggg agagegeeag anucuaagne nggaggueeu gugtyegaue 60
 cacagaauuc qn
                                                              72
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